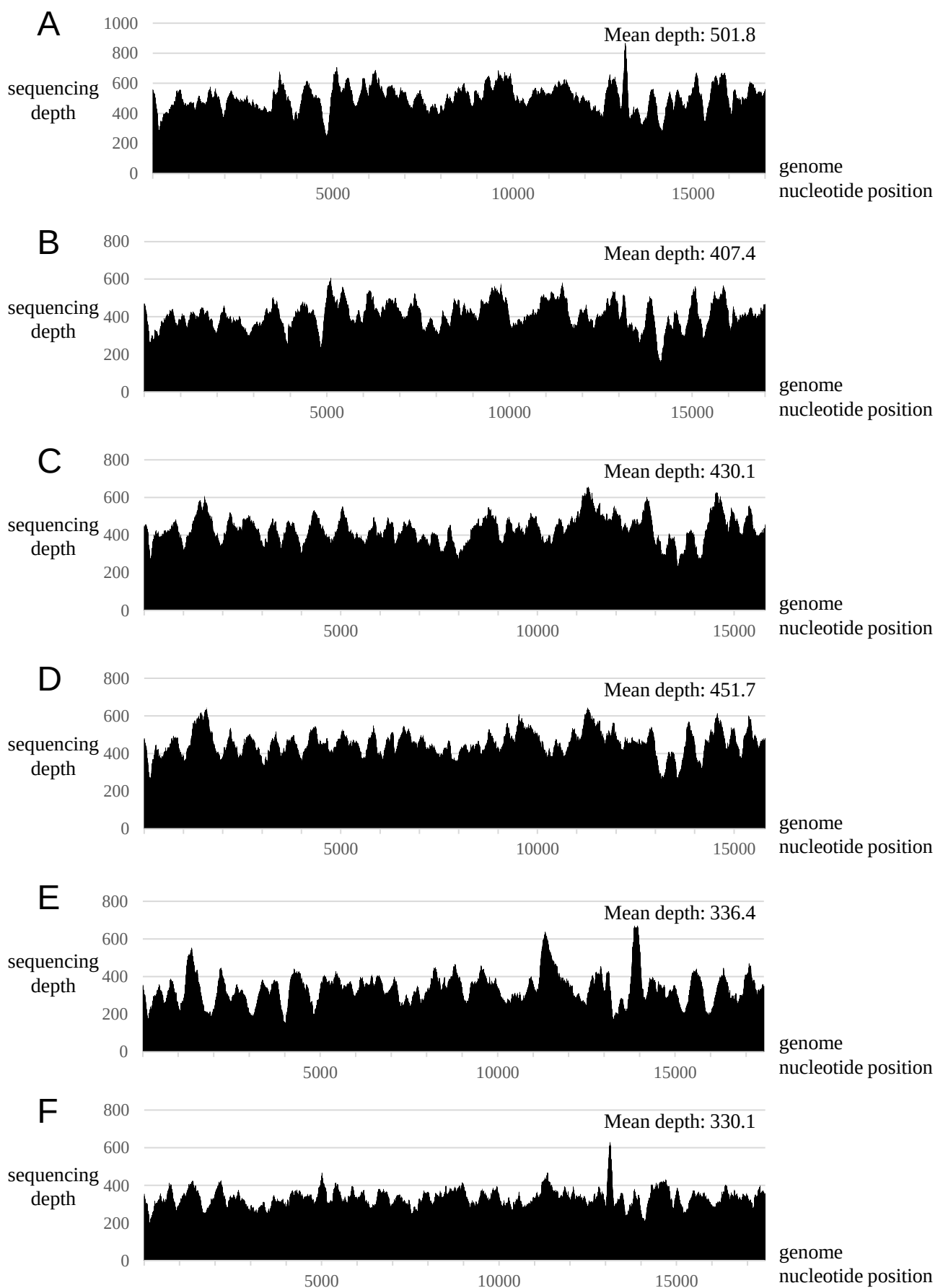




Supplementary Fig. S1 Sequencing depth of reads remapped to mitochondrial genomes of *E. hatanakai* (Eh1) (A), *E. hatanakai* (Eh6) (B), *E. uchidai* (Eu17) (C), *E. uchidai* (Eu36) (D), *E. asanumai* (Ea3) (E), and *E. asanumai* (Ea4) (F). Vertical axis: sequencing depth, horizontal axis: genome nucleotide position.